Run on: October 5, 2005, 11:24:21 ; Search time 168 Seconds
                                        (without alignments)
                                        2115.671 Million cell updates/sec

Title: US-10-616-624-2
Perfect score: 4801
Sequence: 1 MALSWNEIRRKAIEFSKRWE.........QKMTSLLPPEKEIKKSKGKN 919

Scoring table: BLOSUM62
                 Gapop 10.0 , Gapext 0.5

Searched: 2105692 seqs, 386760381 residues

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :   A_Geneseq_16Dec04:*
                 1: geneseqp1980s:*
                 2: geneseqp1990s:*
                 3: geneseqp2000s:*
                 4: geneseqp2001s:*
                 5: geneseqp2002s:*
                 6: geneseqp2003as:*
                 7: geneseqp2003bs:*
                 8: geneseqp2004s:*

                        GenCore version 5.1.6
                 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: October 5, 2005, 11:26:21 ; Search time 47 Seconds
                                        (without alignments)
                                        1881.345 Million cell updates/sec

Title: US-10-616-624-2
Perfect score: 4801
Sequence: 1 MALSWNEIRRKAIEFSKRWE.........QKMTSLLPPEKEIKKSKGKN 919

Scoring table: BLOSUM62
                 Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :   PIR_79:*
                 1: pir1:*
                 2: pir2:*
                 3: pir3:*
                 4: pir4:*

                        GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: October 5, 2005, 11:24:51 ; Search time 183 Seconds
                                             (without alignments)
                                             2571.589 Million cell updates/sec

Title: US-10-616-624-2
Perfect score: 4801
Sequence: 1 MALSWNEIRRKAIEFSKRWE..........QKMTSLLPPEKEIKKSKGKN 919

Scoring table: BLOSUM62
                Gapop 10.0 , Gapext 0.5

Searched: 1612378 seqs, 512079187 residues

Total number of hits satisfying chosen parameters: 1612378

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :   UniProt_03:*
                1: uniprot_sprot:*
                2: uniprot_trembl:*